

Candida attachment to oral epithelium

Vitkov L, Krautgartner WD, Hannig M, Weitgasser R, Stoiber W. Candida attachment to oral epithelium.

Oral Microbiol Immunol 2002: 17: 60-64. © Munksgaard, 2002.

Inflamed oral mucosa biopsies from patients with thrush and high candidal density were observed in a transmission electron microscope (TEM) using ultrahistochemical staining with ruthenium red for glycocalyx visualization. Fimbriae comprising the glycocalyx and enabling yeast adhesion to epithelial cells were clearly visualized by ruthenium red. All internalized portions of the yeast walls were devoid of glycocalyx, indicating that the growing tips of the hyphae mechanically penetrated the host cells. The attachment of *Candida* occurred in two ways: by fimbria-mediated adhesion enabling colonization of the epithelial surface, and by invasion of the superficial epithelial cells via hyphae. As the interaction between adhesin receptors and adhesins stimulates the production of proinflammatory cytokines, *Candida* adhesion itself is assumed to induce mucosal inflammation.

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Key words: adhesins; Candida; fimbriae; oral epithelium; transmission electron microscopy

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Accepted for publication June 28, 2001

The initial stage of epithelial colonization by Candida is thought to be the adherence of the yeast cell to the mucosal surface (16, 22). The ability of a yeast strain to adhere to human epithelial cells has been correlated with virulence in animal models (8, 22). Adhesion takes place by means of long, thin filamentous cell surface appendages termed fimbriae. These form an extracellular coat (glycocalyx). Candida can easily be induced to form fimbriae in vitro by addition of a high concentration of carbohydrates (6, 18, 24). Candidal fimbriae formed in vitro have been visualized in a transmission electron microscope (TEM) by ruthenium red staining (19), similarly to bacterial fimbriae (10). Although it is not possible to maintain such a permanently high concentration of carbohydrates in the oral cavity, other mechanisms could stimulate yeast fimbrial formation in oral Candida-associated diseases. Similarly, bacteria produce glycocalyx in vitro under the influence of high carbohydrate concentrations (5). Recently, electron microscope investigations have demonstrated bacterial attachment to the palatal epithelium in denture stomatitis occurring by fimbria-mediated adhesion (25). Elucidation of the nature of *Candida* attachment to the human oral epithelium in *Candida*-induced diseases will contribute to understanding of the pathogenicity of this yeast.

The aim of this work was to investigate, by electron microscopy, the attachment of *Candida* to the oral epithelium in patients with thrush.

Materials and methods

Thrush, as an exacerbation of previously known stomatitis, was diagnosed by routine oral inspections in six subjects with denture stomatitis and in two spray inhalers (Berodual-Dosier aerosol; Boehringer, Ingelheim, Germany; 3 times a day) with iatrogenic stomatitis of the soft palate. All patients were in the age bracket of 41–71 years (mean: 60 years). The spray inhalers were denture wearers without denture stomatitis and had an intake of 10 mg prednisolon weekly. The criteria suggested by Holmstrup and Axéll (15) were used for diagnosis of thrush. The

study was approved by the local ethics committee, and informed written consent was obtained from all patients. The Candida density was estimated as described previously (27). In addition, samples for microbiological investigation were taken with swabs (Biotest Transportsystem, Biotest AG, Dreieich, Germany) from about 1 cm² of the mucosa, near to the place where the excision was to be made. Sampling took place between 0900 and 1200h, at least 2h after the patient's breakfast. For aerobic culturing, each swab was plated under aerobic conditions on Columbia sheep blood agar (Biotest AG), Columbia CNA agar (Becton Dickinson, Franklin Lakes, NJ) and MacConkey agar (Biotest AG), and incubated at 37°C for 18h. For anaerobic culturing, the swabs were plated on Schaedler agar (Biotest AG) and Schaedler KV agar (Becton Dickinson), and incubated at 37°C for 7 days. Microbiological differentiation was performed as follows.

Viridans streptococci were identified by colony morphology, alpha-haemolysis, and use of a commercially available optochin test. Nonhaemolytic streptococci were identified by colony morphology, lack of haemolysis and the optochin test. Enterococci were identified by colony morphology and antibiotic susceptibility. Staphylococci were identified by colony morphology, colony colour, and testing of coagulase activity with Staphaurex (Abbott Laboratories, Abbott Park, IL). Hemophilus spp. were identified by colony morphology, gram staining, and biochemical identification procedures API NH (Biomerieux SA, Marcy l'Etoile, France). Neisseria spp. were identified by colony morphology, gram staining and positive oxidase reaction. Gram-negative bacilli (Escherichia coli, Klebsiella, Enterobacter, etc.) as well as Lactobacillus casei were identified by use of a biochemical identification system (Crystal System, Becton Dickinson, Franklin Lakes, NJ).

For semiquantitative estimation of the bacterial density, the following counting procedure was used.

- Negative bacterial growth was characterized by 0-4 colonies per agar plate.
- Low bacterial density was defined by the growth of at least five colonies per agar plate.
- Moderate bacterial density was defined by growth over half of the agar plate.
- High bacterial density was defined by confluent bacterial growth.

Biopsies (approx. $5 \times 8 \text{ mm}$) were taken from the inflamed mucosa of each patient under regional block anaesthesia. Each biopsy was washed in 50 ml saline by gently swaying for 20 s and

subsequently divided into two equal parts. The specimens were washed to remove all substances in contact with but not attached to the epithelial surface. One sample from each patient was fixed with 1.2% glutaraldehyde (buffered at pH 6.5 with 0.1 M sodium cacodylate) for 2h at 4°C. Postfixation of these control samples was performed with 1% osmium tetroxide (buffered at pH 6.5 with 0.1 M sodium cacodylate) for 2h at 4°C. The other sample from each patient was fixed and postfixed by the fixatives mentioned above with addition of 0.05% ruthenium red in each case. All specimens were routinely embedded in Epon 812. Ultrathin sections were cut on an ultramicrotome (Reichert Ultracut S, Optische Werke C. Reichert, Vienna, Austria), contrasted with LKB 2168 Ultrostainer (LKB Produkter AB, Bromma, Sweden) and examined in a transmission electron microscope LEO EM 910 (LEO Elektronenmikroskopie Ltd. Oberkochen, Germany) operating at 80

Results

Results of the microbiological examination are summarized in Table 1. The predominant yeast species found in the samples was *Candida albicans*. Certain differences concerning the bacterial flora were detected between individuals. The normal oral bacteria were represented by viridans streptococci and *Neisseria* spp. *Staphylococcus epidermidis* was found in 50% of the patients, and anaerobes were found in 12%.

Electron-microscopically, numerous yeasts were identified intra- and extracellularly (Fig. 1). Different Candida cell numbers were observed between and within individual samples. A fairly uniform number of circular or ovoid transverse sections of the fungi were present in the ultra-thin sections, making it difficult to distinguish between sections through spherical/ovoid blastospores and tubular hyphae. The cell walls and cytoplasmic contents were similar in all circular, ovoid and tubular yeast sections. The yeast was predominantly located on and/or within desquamated and loose epithelial cells (Fig. 1).

The samples fixed in the absence of ruthenium red showed no (Fig. 2) or a rather faint layer exterior to the cell wall (Fig. 3). However, staining with ruthenium red revealed the existence of a thick layer of fibrils with high electron

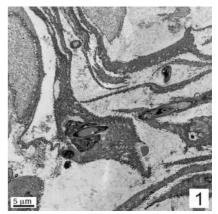


Fig. 1. Control staining, showing loose palatal epithelium with extra- and intracellular yeasts.

Table 1.	Results	of the	microbiological	investigation:	yeast	and	bacterial	species	and	density
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Patient	Age (yr)	Sex	Candida spp.	Viridans streptococci	Neisseria spp.	Enteroccocus faecalis	Staphylococcus epidermidis	Escherichia coli	Enterobacter aerogenes	Klebsiella oxytoca	Morganella marganii	Bacteroides spp.
1	49	m	alb+			+++			+++			
			spp ++									
2	61	m	alb +++	++			+++	+++		+++		
3	71	m	alb +++	++		++	++					
4	41	f	alb+	++	++							
			spp ++									
5	60	m	alb +++	++			+	+++				
6	64	f	alb +++			+						
7	68	m	alb +++	++		++	++					+
8	66	m	alb +++	+++	++					++	++	

⁺⁼ low yeast or bacterial density; ++= moderate density; +++= high density; alb = Candida albicans; spp = Candida non-albicans species.

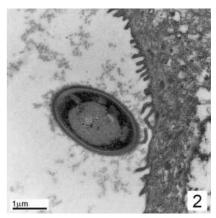


Fig. 2. Control staining, showing absence of a layer exterior to the yeast cell wall, and thickening of the epithelial cytoplasm around the yeast.

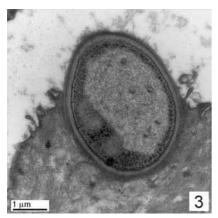


Fig. 3. Control staining, showing a faint layer exterior to the yeast cell wall outside the epithelial cell.

density exterior to the cell wall (Figs 4, 8 and 9). The TEM analysis showed yeast adhesion to the epithelium and internalization in all biopsy samples. The fimbriae were not always easily distinguishable (Figs 4 and 7) because of their high electron density, but in some cases they were clearly distinguishable on the eukaryotic cell membrane (Figs 5, 6 and 8). All parts of the internalized yeast walls were devoid of glycocalyx (Fig. 6).

In cases where hyphal tip growth towards the loosened epithelial cell was observed, a deformation of the epithelial membrane and cytoskeleton and the formation of an epithelial cell invagination around the depressing hyphal tip were seen (Figs 7 and 9).

All areas where bacteria were observed were located superficially scattered on the epithelial surface. High ultrastructural diversity (cocci, coccus-

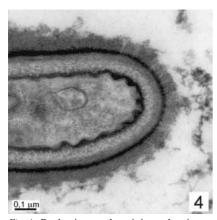


Fig. 4. Ruthenium red staining, showing a large yeast glycocalyx with very electrondense contrasting exterior to the cell wall.

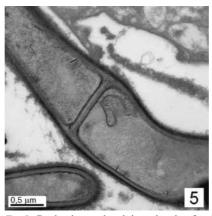


Fig. 5. Ruthenium red staining, showing formation of a dividing wall between two true hyphae.

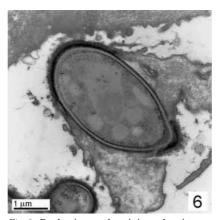


Fig. 6. Ruthenium red staining, showing an epithelial cell penetrated by a yeast, with glycocalyx formation on the yeast cell wall only outside the epithelial cell.

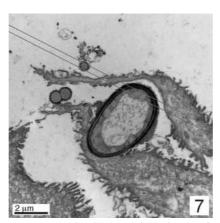


Fig. 7. Ruthenium red staining of a longitudinal section through an epithelial cell invagination containing a growing hypha. The epithelial cell surface is distinctly deformed. The two lines mark the boundary of a cross-section as demonstrated below in Fig. 9. Diplococcus or dividing cocci can be seen adhering to the epithelial cell membrane.

like and rod-shaped bacteria) was observed in the bacteria adhering to the outer surface of the epithelial cell membranes (Fig. 7) or to other bacteria (Fig. 9). In addition, a certain morphological similarity between bacterial (Fig. 10) and yeast (Fig. 8) glycocalyces, both consisting of fimbriae, was evident.

Discussion

The present study provides, for the first time, ultrastructural evidence of fimbriae-mediated *Candida* adhesion to human oral epithelium in a *Candida*-in-

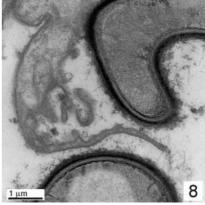


Fig. 8. Ruthenium red staining, showing separate fimbriae adhering to the epithelial cell membrane.

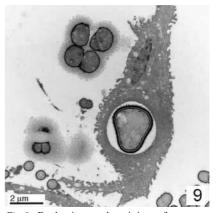


Fig. 9. Ruthenium red staining of a crosssection through a shallow epithelial cell invagination containing a growing hypha. The very electron-dense glycocalyx contrasted by ruthenium red provides unequivocal evidence of the extracellular position of the fungus, since ruthenium red is unable to penetrate the epithelial cell membrane (8). The separate fimbriae adhering to the epithelial cell membrane are clearly distinguishable. Bacteria adhered by glycocalyces with different largeness.

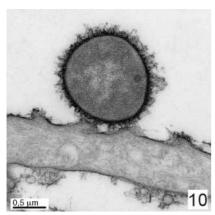


Fig. 10. Ruthenium red staining, showing a bacterium adhering by fimbriae to the epithelial cell membrane.

duced disease. The adhesins recognize and bind to particular receptors expressed on the eukaryotic cell membrane. Morphologically, the *Candida* adhesins are integrated into fimbriae. The major structural unit of the *C. albicans* fimbriae consists of 80–85% carbohydrate (primarily D-mannose) and 10–15% protein, so that the fimbriae are biochemically analogous to the bacterial fimbriae (29). Some morphological similarity between the fimbriae of bacteria and *Candida* is also indicated in the present TEM micrographs.

As carbohydrates are not stainable by routine contrasting substances, glycocalyx cannot be easily monitored in TEM, unless special staining techniques are used. The ability of cationic ruthenium red to enhance staining of acidic glycosylated structures at the ultrastructural level is widely used for the visualization of such structures (9, 10,14, 25). The extracellular fungi stained by ruthenium red displayed a thick glycocalyx, which was seen in some cases in control staining as a faint coating, similarly to the floccular layer described in oral candidiasis (21) and above the denture plaque (26). Electron-microscopically, the glycocalyx of the oral yeasts showed the same base characteristics as the glycocalyx of Candida grown in vitro with addition of 0.5 M sucrose (18) or 0.5 M galactose (6). The partial visualization of Candida glycocalyx in nonruthenium-red-stained specimens could result from staining the proteinaceous part of the fimbriae or proteins adsorbed on them (21).

Epithelial cells cannot be penetrated by blastospores, but only by the growing hyphal tips (16). The attachment via glycocalyx to some surfaces provides anchorage for the hyphae and allows growth through the epithelial tissue. Consequently, penetration can occur only if anchorage by adhesion has taken place in advance.

An adhesin constituent, the supernatant of the mannoprotein fraction (MP-F2), elicits a greatly delayed type hypersensitivity reaction, Th1 and Th2 cytokine production and anticandidal protection in mice (20). Another glycocalyx constituent, beta-1,2-linked oligomannoside residues of C. albicans, acts as an adhesin for macrophages and stimulates these cells to undergo cytokine production (11). C. albicans mannan stimulates the production of IL-2, IL-4 and gamma interferon by mouse splenocytes (17). In addition, the mannose receptors of macrophages are involved in cytokine IL-1B, IL-6 and granulocyte-macrophage colony-stimulating factor responses (28). As the interaction between adhesin receptors and adhesins stimulates the production of proinflammatory cytokines, the fimbria-mediated adhesion of Candida could consequently be regarded as a primary inducer of the Candida-induced inflammation, which is further enhanced by Candida toxins (4, 7) and enzymes (4, 19).

C. albicans has the ability to adhere in vitro to the host cell surface receptors via a number of adhesins (30). The fimbriae of Candida possibly interact with the glycosphingolipid receptors via the carbohydrate portion of the receptors, displayed on the surface of human buccal epithelial cells (30). In addition, purified adhesins of C. albicans bind to glycolipids carrying the H blood group antigen (3). Among possible candidates for Candida receptors on epithelial cells are the ABO and Lewis blood group antigens (overall H and Lewis blood group antigens), all of which posses residues of L-fucose (3, 23). An association was demonstrated between blood group O, with nonsecretion of blood group antigens, and Candida carriage (1, 2). It is very likely that the salivary blood group antigens bind to the Candida adhesins and hinder the adhesion of the yeasts to the cytoplasmic membrane surfaces. Therefore, host susceptibility to the opportunistic pathogen Candida might largely be determined by the expression of adhesin receptors (blood group antigens) on the cytoplasmic membrane surfaces and particularly by nonsecretion of blood group antigens in saliva.

Antiserum against adhesins protects against disseminated candidiasis in animal models (12), and a vaccine against adhesin in animal models induces production of a monoclonal antibody protecting against candidiasis (13). It should be possible to develop an immunization protocol for *Candida* adhesins that leads to memory B-cell responses and long-term immunity against various forms of candidiasis (13).

The ultrastructural evidence presented for fimbria-mediated adhesion in thrush highlights the role of adhesion as a triggering factor in *Candida*-induced oral pathology.

Acknowledgments

The authors thank Mrs Adelina Vitkov for secretarial assistance. This work was supported by the Medical Association of Salzburg, Austria and the Medical Research Society of Salzburg, Austria.

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